

S2_Table: Study design for transcriptome profiling of rabbit ear wound responses to *P.aeruginosa* infection.

Group	Biological Repeats	CFU Inoculated per Wound	Number of Rabbits ^a used in the study						Sub-totals
			^b PWD0	PWD3 (2 h post-infection)	PWD3 (6h post-infection)	PWD4 (24 h post-infection)	PWD8 (5 d post-infection)	PWD12 (9 d post-infection)	
1	A	0	3	3	3	3	3	3	18
2		1×10 ⁶	0	5	5	5	4	4	23
3	B	0	3	3	3	3	3	3	18
4		1×10 ⁶	0	5	5	5	4	4	23
5	C	0	-	-	-	-	3	-	3
6		1×10 ⁶	-	-	-	-	4	-	4
		Total Rabbits used in this study							89

^a 12 (6 mm) full thickness dermal excisional wounds were produced per rabbit (6 wounds per ear). All wounds in control and experimental animals were treated the same.

^bPWD is Post Wounding Day